

**In the Specification**

Please enter the enclosed SEQUENCE LISTING into the specification.

Please amend the paragraph beginning at line 18 of page 8 as follows:

Figure 1 provides a CLUSTAL X (version 1.8) multiple sequence alignment of the cleavage spanning regions of select MMP proenzymes (SEQ ID NOs:2-10). Figure 1A provides an alignment that highlights conserved residues where an '\*' indicates complete identity among the sequences, a ':' indicates 7/9 conserved positions, and a '.' indicates greater than 80% identical positions with mostly conserved substitutions. Figure 1B indicates the positions of heterogeneity in bold.